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IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In Re Application of:

LAI et al.

Confirmation No.: 4378

Serial No. 09/846,430

Group Art Unit: 1642

Filing Date: April 30, 2001

Examiner: Tara Washington

Title: METHODS AND COMPOSITIONS FOR POLYNUCLEOTIDE
ANALYSIS USING GENERIC CAPTURE SEQUENCES

TRANSMITTAL LETTER

Assistant Commissioner for Patents
Washington, D.C. 20231

Sir:

Transmitted herewith for filing is an Information Disclosure Statement, including a Form PTO-1449 and copies of the cited references. It is believed that no fee is due.

The Commissioner is hereby authorized to charge any fees under 37 C.F.R. §§ 1.16, 1.17 and 1.21 which may be required by this paper, or to credit any overpayment, to Deposit Account No. 18-1648.

Respectfully submitted,

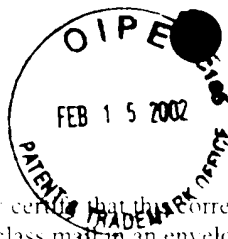
Date:

January 29/2002

By:

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**INFORMATION DISCLOSURE STATEMENT
UNDER 37 C.F.R. § 1.97**

Assistant Commissioner for Patents
Washington, D.C. 20231

Sir:

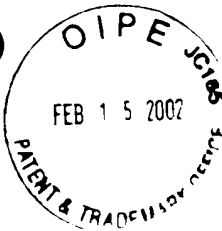
The information listed below may be material to the examination of the above-identified application. Copies of the information and completed PTO-1449 forms are submitted herewith. The check for \$180 to cover the fee is attached. The Examiner is respectfully requested to make this information of official record in the application. The information includes:

United States Patent No. 6,287,778 B1 issued Sept. 11, 2001, to Huang et al.;

Brenner et al., "Gene Expression Analysis by Massively Parallel Signature Sequencing (MPSS) on Microbead Arrays," *Nature Biotechnology* 18:630-634 (2000);

Chen et al., "A Microsphere-Based Assay for Multiplexed Single Nucleotide Polymorphism Analysis Using Single Base Chain Extension," *Genome Research* 10:549-557 (2000);

Favis et al., "Universal DNA Array Detection of Small Insertions and Deletions in *BRC11* and *BRC12*," *Nature Biotechnology* 18:561-564 (2000);



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Gerry et al., "Universal DNA Microarray Method for Multiplex Detection of Low Abundance Point Mutations," *J. Mol. Biol.* 292:251-262 (1991);

Iannone et al., "Multiplexed Single Nucleotide Polymorphism Genotyping by Oligonucleotide Ligation and Flow Cytometry," *Cytometry* 39:139-140 (2000);

Myakishev et al., "High-Throughput SNP Genotyping by Allele-Specific PCR with Universal Energy-Transfer-Labeled Primers," *Genome Research* 11:163-169 (2001); and

Tillib and Mirzabekov, "Advances in the Analysis of DNA Sequence Variations Using Oligonucleotide Microchip Technology," *Analytical Biotechnology* 12:53-58 (2001).

This Information Disclosure Statement under 37 CFR § 1.97 is not to be construed as a representation that: (i) a complete search has been made; (ii) additional information material to the examination of this application does not exist; (iii) the information, protocols, results and the like reported by third parties are accurate or enabling; or (iv) the above information constitutes prior art to the subject invention.

Respectfully submitted,

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